## AMENDMENTS TO THE CLAIMS

- 1. (Currently Amended) A modified glucose dehydrogenase having composition comprising water-soluble *Acinetobacter calcoaceticus* pyrroloquinoline quinone (PQQGDH) as a coenzyme, wherein one or more amino acid residues in a region of 349-377 amino acid amino acid Thr366 Thr342 and/or amino acid Asp167 Asp143 of SEQ ID NO: 1 of water-soluble the PQQGDH derived from *Acinetobacter calcoaceticus* is are replaced with other amino acid residues, and wherein said PQQGDH has an inhibition constant (Ksi) of 200 mM or more.
- 2. (Withdrawn) A modified glucose dehydrogenase having pyrroloquinoline quinone as a coenzyme, wherein Met365 of the amino acid sequence defined in SEQ ID NO:1 is replaced with another amino acid, and has a Ksi value of 200 mM or more.
- 3. (Withdrawn) A modified glucose dehydrogenase having pyrroloquinoline quinone as a coenzyme, wherein Met365 of the amino acid sequence defined in SEQ ID NO:1 is replaced with tryptophan or phenylalanine.
- 4. (Currently Amended) A <u>The modified glucose dehydrogenase according to claim 1</u>, having pyrroloquinoline quinone as a coenzyme wherein <u>Thr366 Thr342</u> of the amino acid sequence defined in SEQ ID NO: 1 of <u>PQQGDH</u> is replaced with another amino acid, and has a <u>Ksi value of 200 mM or more</u>.
- 5. (Currently Amended) A <u>The modified glucose dehydrogenase according to claim 1</u>, having pyrroloquinoline quinone as a coenzyme wherein <u>Thr366 Thr342</u> of the amino acid sequence defined in SEQ ID NO: 1 <u>of PQQGDH</u> is replaced with aspartic acid, lysine, isoleucine, or <u>asparagines asparagine</u>.

6. (Withdrawn) A modified glucose dehydrogenase having pyrroloquinoline quinone as a coenzyme wherein Tyr367 of the amino acid sequence defined in SEQ ID NO: 1 is replaced with another amino acid, and has a Ksi value of 200 mM or more.

- 7. (Withdrawn) A modified glucose dehydrogenase having pyrroloquinoline quinone as a coenzyme wherein Tyr367 of the amino acid sequence defined in SEQ ID NO: 1 is replaced with aspartic acid.
- 8. (Withdrawn) A modified glucose dehydrogenase having pyrroloquinoline quinone as a coenzyme wherein Ile368 of the amino acid sequence defined in SEQ ID NO: 1 is replaced with another amino acid, and has a Ksi value of 200 mM or more.
- 9. (Withdrawn) A modified glucose dehydrogenase having pyrroloquinoline quinone as a coenzyme wherein Ile368 of the amino acid sequence defined in SEQ ID NO: 1 is replaced with asparagine.
- 10. (Withdrawn) A modified glucose dehydrogenase having pyrroloquinoline quinone as a coenzyme wherein Cys369 of the amino acid sequence defined in SEQ ID NO: 1 is replaced with another amino acid and has a Ksi value of 200 mM or more.
- 11. (Withdrawn) A modified glucose dehydrogenase having pyrroloquinoline quinone as a coenzyme wherein Cys369 of the amino acid sequence defined in SEQ ID NO: 1 is replaced with arginine.
- 12. (Withdrawn) A modified glucose dehydrogenase having pyrroloquinoline quinone as a coenzyme wherein Ala374 of the amino acid sequence defined in SEQ ID NO: 1 is replaced with another amino acid, and has a Ksi value of 200 mM or more.

13. (Withdrawn) A modified glucose dehydrogenase having pyrroloquinoline quinone as a coenzyme wherein Ala374 of the amino acid sequence defined in SEQ ID NO: 1 is replaced with proline.

## 14. - 15. (Cancelled)

- 16. (Currently Amended) A <u>The</u> modified glucose dehydrogenase <u>according to claim 1</u>, having pyrroloquinoline quinone as a coenzyme wherein an amino acid residue selected from the group consisting of Met365, Thr366, <u>Thr342</u> <u>Tyr367, Ile368, Cys369</u>, and Ala374 of the amino acid sequence defined in SEQ ID NO:1 <u>of PQQGDH</u> is replaced with another amino acid and wherein <u>Asp167</u> <u>Asp143 of SEQ ID NO:1</u> is replaced with glutamic acid.
- 17. (Currently Amended) A <u>The</u> modified glucose dehydrogenase <u>according to claim 1</u>, having pyrroloquinoline quinone as a coenzyme wherein <u>Thr366 Thr342</u> of the amino acid sequence defined in SEQ ID NO:1 <u>of PQQGDH</u> is replaced with aspartic acid, lysine, isoleucine, or asparagine, and wherein <u>Asp167 Asp143 of SEQ ID NO:1</u> is replaced with glutamic acid.
- 18. (Withdrawn) A glucose dehydrogenase having pyrroloquinoline quinone as a coenzyme comprising the following amino acid sequence:

Cys Gly Glu Xaa Thr Tyr Ile (SEQ ID NO:3)

wherein Xaa is Met or Trp.

19. (Withdrawn) A glucose dehydrogenase having pyrroloquinoline quinone as a coenzyme comprising the following amino acid sequence:

Gly Glu Met Xaa Tyr Ile Cys (SEQ ID NO:4)

wherein Xaa is Asp, Lys, Ile or Asn.

20. (Withdrawn) A glucose dehydrogenase having pyrroloquinoline quinone as a coenzyme comprising the following amino acid sequence:

Glu Met Thr Asp Ile Cys Trp (SEQ ID NO:5).

- 21. (Withdrawn) A glucose dehydrogenase having pyrroloquinoline quinone as a coenzyme comprising the following amino acid sequence:

  Met Thr Tyr Asp Cys Trp (SEQ ID NO:6).
- 22. (Withdrawn) A glucose dehydrogenase having pyrroloquinoline quinone as a coenzyme comprising the following amino acid sequence:

  Thr Tyr Ile Arg Trp Pro Thr (SEQ ID NO:7).
- 23. (Withdrawn) A glucose dehydrogenase having pyrroloquinoline quinone as a coenzyme comprising the following amino acid sequence:

  Pro Thr Val Pro Pro Ser (SEQ ID NO:8).
- 24. (Withdrawn) A gene encoding a modified glucose dehydrogenase as claimed in claim 1.
  - 25. (Withdrawn) A vector comprising the gene as claimed in claim 24.
  - 26. (Withdrawn) A transformant comprising the gene as claimed in claim 24.
- 27. (Withdrawn) A transformant as claimed in claim 26, wherein the gene as claimed in claim 24 is integrated in its chromosome.

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28. (Withdrawn) A method for preparing a water-soluble PQQGDH, comprising culturing the transformant as claimed in claim 27 and preparing water-soluble fraction from the cells of the transformant.

- 29. (Currently Amended) A glucose assay kit comprising the modified glucose dehydrogenase as claimed in according to claim 1.
- 30. (Currently Amended) A glucose sensor comprising the modified glucose dehydrogenase as claimed in according to claim 1.